

SEQUENCE LISTING

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<120> Method for Triggering Apoptosis in Cells

<130> apoptosis

<140> 09/701,618
<141> 2001-02-14

<150> PCT/DE99/01684
<151> 1999-06-02

<150> DE 198 24 811.3
<151> 1998-06-03

<160> 10

<170> PatentIn Ver. 2.1

<210> 1
<211> 1156
<212> DNA
<213> Homo sapiens

<220>
<221> CDS
<222> (118)..(540)
<223> Human C1D cDNA

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tgtgaggcaa gagtacctat agaaccggaa ggagggtgag gagcagagct ggccata 117
atg gca ggt gaa gaa att aat gaa gac tat cca gta gaa att cac gag 165
Met Ala Gly Glu Glu Ile Asn Glu Asp Tyr Pro Val Glu Ile His Glu
1 5 10 15
tat ttg tca gcg ttt gag aat tcc att ggt gct gtg gat gag atg ctg 213
Tyr Leu Ser Ala Phe Glu Asn Ser Ile Gly Ala Val Asp Glu Met Leu
20 25 30

aag acc atg atg tct gtt tct aga aat gag ttg ttg cag aag ttg gat	261		
Lys Thr Met Met Ser Val Ser Arg Asn Glu Leu Leu Gln Lys Leu Asp			
35	40	45	
cca ctt gaa caa gca aaa gtg gat ttg gtt tct gca tac aca tta aat	309		
Pro Leu Glu Gln Ala Lys Val Asp Leu Val Ser Ala Tyr Thr Leu Asn			
50	55	60	
tca atg ttt tgg gtt tat ttg gca acc caa gga gtt aat cct aag gaa	357		
Ser Met Phe Trp Val Tyr Leu Ala Thr Gln Gly Val Asn Pro Lys Glu			
65	70	75	80
cat cca gta aaa cag gaa ttg gaa aga atc aga gta tat atg aac aga	405		
His Pro Val Lys Gln Glu Leu Glu Arg Ile Arg Val Tyr Met Asn Arg			
85	90	95	
gtc aag gaa ata aca gac aag aaa aag gct ggc aag ctg gac aga ggt	453		
Val Lys Glu Ile Thr Asp Lys Lys Ala Gly Lys Leu Asp Arg Gly			
100	105	110	
gca gct tca aga ttt gta aaa aat gcc ctc tgg gaa cca aaa tcg aaa	501		
Ala Ala Ser Arg Phe Val Lys Asn Ala Leu Trp Glu Pro Lys Ser Lys			
115	120	125	
aat gca tca aaa gtt gcc aat aaa gga aaa agt aaa agt taacttttg	550		
Asn Ala Ser Lys Val Ala Asn Lys Gly Lys Ser Lys Ser			
130	135	140	
gttttcatgt acacatattc aaaaagtaca ttaatatgtt atcacagtaa tatgtaaagc	610		
taaatacttc ctctccaaag atcattatct ttattgatta gcactgagga tttaacatt	670		
gtgatatat attatattat aatttaccat ctcttgatga gactcttatt tctttatata	730		
ggtcagtctt gcaagtacca ttttataagc agctgtgaaa tttaagtgaa atgttcttg	790		
taaacatttg tactattta aatgaataat gaccttatga agtatgctat ctgtaggctg	850		
aaattatagg tacatctgtt ttcactatat gatattaaga aagcgtgaat gacttaatg	910		
ttcattttt tctgtataga tactttatca tgtttcatg attttaggaa ttactgcttt	970		
gttcatattc aaagtgtgaa actaaaagtt tatgggtgta cttaattct tggcatgttg	1030		
cctctatgtc ccattaaaaaaa taaaatacat tctcattaac ttttagatggg aaataagggtt	1090		
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gtaaaa

1156

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<211> 141
<212> PRT
<213> Homo sapiens

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Tyr Leu Ser Ala Phe Glu Asn Ser Ile Gly Ala Val Asp Glu Met Leu
20 25 30
Lys Thr Met Met Ser Val Ser Arg Asn Glu Leu Leu Gln Lys Leu Asp
35 40 45
Pro Leu Glu Gln Ala Lys Val Asp Leu Val Ser Ala Tyr Thr Leu Asn
50 55 60
Ser Met Phe Trp Val Tyr Leu Ala Thr Gln Gly Val Asn Pro Lys Glu
65 70 75 80
His Pro Val Lys Gln Glu Leu Glu Arg Ile Arg Val Tyr Met Asn Arg
85 90 95
Val Lys Glu Ile Thr Asp Lys Lys Ala Gly Lys Leu Asp Arg Gly
100 105 110
Ala Ala Ser Arg Phe Val Lys Asn Ala Leu Trp Glu Pro Lys Ser Lys
115 120 125
Asn Ala Ser Lys Val Ala Asn Lys Gly Lys Ser Lys Ser
130 135 140

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<212> DNA
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<220>
<221> CDS
<222> (78)..(500)
<223> Mouse C1D cDNA

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 Met Ala Gly Glu Glu Met Asn Glu Asp Tyr Pro
 1 5 10
 gta gaa att cac gag tct tta aca gcc ctg gag agc tcc ctg ggt gct 158
 Val Glu Ile His Glu Ser Leu Thr Ala Leu Glu Ser Ser Leu Gly Ala
 15 20 25
 gtg gac gac atg ctg aag acc atg atg gct gtt tct aga aac gag ttg 206
 Val Asp Asp Met Leu Lys Thr Met Met Ala Val Ser Arg Asn Glu Leu
 30 35 40
 ttg cag aag ttg gac cca ttg gaa caa gca aag gtg gat tta gtt tct 254
 Leu Gln Lys Leu Asp Pro Leu Glu Gln Ala Lys Val Asp Leu Val Ser
 45 50 55
 gca tac acc tta aat tca atg ttt tgg gtt tat ttg gca act caa gga 302
 Ala Tyr Thr Leu Asn Ser Met Phe Trp Val Tyr Leu Ala Thr Gln Gly
 60 65 70 75
 gtt aat ccc aaa gag cat cca gtg aag cag gaa ctg gaa aga atc aga 350
 Val Asn Pro Lys Glu His Pro Val Lys Gln Glu Leu Glu Arg Ile Arg
 80 85 90
 gtc tac atg aac aga gtt aaa gaa ata aca gac aag aag gct gcc 398
 Val Tyr Met Asn Arg Val Lys Glu Ile Thr Asp Lys Lys Ala Ala
 95 100 105
 aag ctg gac aga ggt gct tcg aga ttt gtc aag aag gca ctc tgg 446
 Lys Leu Asp Arg Gly Ala Ala Ser Arg Phe Val Lys Lys Ala Leu Trp
 110 115 120
 gaa ccc aaa cga aaa agc aca cca aaa gtg gct aat aaa ggg aaa agc 494
 Glu Pro Lys Arg Lys Ser Thr Pro Lys Val Ala Asn Lys Gly Lys Ser
 125 130 135
 aaa cac taatcttttg gttttgatgt acatgttttc aaaaagtaca tccttttaa 550
 Lys His
 140
 tcagtttaca atgttagttat gtgaccatgt ggtgtttaaa tggattcctt ttggaaattca 610
 tgtataaaatt tacacattac atttgtgata ctgaatctt ttttgctga gaaagattaa 670

gttgtcttg ttgatttca tataaagcat catgatgtgt ttaatattgt aagatattct 730
ataaggcgtt gtgaaatcca aatgttctct gtaaacattt gtagtgtttg aaatgaacaa 790
tgatattatg aagtgtgcta tctgttagacc tcgaggtgta aggacatttgg ttttcagtaa 850
tgatgagaaa tacagtgact taaataccca ctctgttct gttcagtttgc ttcaacatgt 910
ttcgtgattt tttttttttt tttagtaatt ctgtcttgat attcaaagtc aaaattgaaa 970
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tgatggaaaaa 1040

<210> 4
<211> 141
<212> PRT
<213> Mus musculus

<400> 4
Met Ala Gly Glu Glu Met Asn Glu Asp Tyr Pro Val Glu Ile His Glu
1 5 10 15

Ser Leu Thr Ala Leu Glu Ser Ser Leu Gly Ala Val Asp Asp Met Leu
20 25 30

Lys Thr Met Met Ala Val Ser Arg Asn Glu Leu Leu Gln Lys Leu Asp
35 40 45

Pro Leu Glu Gln Ala Lys Val Asp Leu Val Ser Ala Tyr Thr Leu Asn
50 55 60

Ser Met Phe Trp Val Tyr Leu Ala Thr Gln Gly Val Asn Pro Lys Glu
65 70 75 80

His Pro Val Lys Gln Glu Leu Glu Arg Ile Arg Val Tyr Met Asn Arg
85 90 95

Val Lys Glu Ile Thr Asp Lys Lys Ala Ala Lys Leu Asp Arg Gly
100 105 110

Ala Ala Ser Arg Phe Val Lys Lys Ala Leu Trp Glu Pro Lys Arg Lys
115 120 125

Ser Thr Pro Lys Val Ala Asn Lys Gly Lys Ser Lys His
130 135 140

<210> 5
<211> 38
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Forward PCR
Primer for Human cDNA

<400> 5
gggttaccat ggcaggtgaa gaaattaatg aagactat 38

<210> 6
<211> 38
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Reverse PCR
Primer for Human cDNA

<400> 6
gggtcgactt aacttttact ttttccttta ttggcaac 38

<210> 7
<211> 38
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Forward PCR
Primer for Mouse cDNA

<400> 7
gggttaccat ggcaggtgaa gaaatgaatg aagattat 38

<210> 8
<211> 38
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Reverse PCR

Primer for Mouse cDNA

<400> 8 gggtcgacgt gtttgctttt ccctttattta gccacttt 38

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<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Forward PCR
Primer for Enhanced Green Fluorescent Protein

<400> 9 gggtcgacat ggtgagcaag ggcgaggagc tgttc 35

<210> 10
<211> 35
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Reverse PCR
Primer for Enhanced Green Fluorescent Protein

<400> 10 ccaagctttg gaattctaga gtcgcggccg ctta 35